

RAW SEQUENCE LISTING
 ENTRY: 17252000X1613355

File: C:\CRF3\Outfold\Vsrl613355.raw
 (XREF) 17252000X1613355.raw

(1) INFORMATION FOR SEQ ID NO: 1:
 (a) LENGTH: 10
 (b) TYPE: AMINO ACID
 (c) STRANDEDNESS: SINGLE
 (d) TOPOLOGY: LINEAR
 (e) FUNCTIONAL TYPE: PEPTIDE IN SEQ ID NO: 1
 (f) HYDROPHETICAL NO
 (g) ANTI-SENSE
 (v) FRAGMENT TYPE:
 (i) ORIGINAL SOURCE:
 (ii) ORGANISM:
 (iii) STRAIN:
 (iv) INDIVIDUAL ISOLATE:
 (v) EXPERIMENTAL STAGE:
 (vi) HARVESTED:
 (vii) TISSUE TYPE:
 (viii) CELL TYPE:
 (ix) CELL LINE:
 (x) ORGANELLE:
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
 N L G E H P V C D S I D I W V
 (2) INFORMATION FOR SEQ ID NO: 2:
 (a) LENGTH: 10
 (b) TYPE: AMINO ACID
 (c) STRANDEDNESS: SINGLE
 (d) TOPOLOGY: LINEAR
 (e) FUNCTIONAL TYPE: PEPTIDE IN SEQ ID NO: 2
 (f) HYDROPHETICAL NO
 (g) ANTI-SENSE
 (v) FRAGMENT TYPE:
 (i) ORIGINAL SOURCE:
 (ii) ORGANISM:
 (iii) STRAIN:
 (iv) INDIVIDUAL ISOLATE:
 (v) EXPERIMENTAL STAGE:
 (vi) HARVESTED:
 (vii) TISSUE TYPE:
 (viii) CELL TYPE:
 (ix) CELL LINE:
 (x) ORGANELLE:
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 N L G E H P V C D S I D I W V

invalid

delete

invalid

For 1.822 of
 Sequence Rules,
 do not show
 amino or
 Carboxy groups

1) delete all
 dashes;

2) Number the
 amino acids
 under every 5
 amino acids -
 do not use
 TAB codes,
 between amino
 acid nos.

Per Sequence Rules,
 use three-letter
amino acids

PAW SEQUENCE LISTING
 EVENT: A:118A118A US/09/813,355

Input File: A:\Nlpp8seq
 Output File: N:\CRF3\07252000\1613355.14W

W--> 97 (v) FRAGMENT TYPE:
 (1) ORIGINAL SOURCE:
 (2) ORGANISM:
 (3) STRAIN:
 (4) INDIVIDUAL IDENTIFIER:
 (5) DEVELOPMENTAL STAGE:
 (6) HOST TYPE:
 (7) TISSUE TYPE:
 (8) CELL TYPE:
 (9) CELL LINE:
 (I) ORGANEILE:
 C--> 107 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
 C--> 109 N L G E H P V C D S
 E--> 110 INFORMATION FOR SEQ ID NO: 4:
 C--> 112 (2) SEQUENCE CHARACTERISTICS:
 (1) LENGTH: 3
 (2) TYPE: AMINO ACID
 (3) STRAN: FORWARD
 (4) SOURCE: BENT
 (5) ORGANISM: HUMAN
 (6) TISSUE TYPE: PLACENTA IN SEQ ID NO: 1
 (7) DEVELOPMENTAL STAGE: N
 (8) HOST TYPE:
 (9) CELL TYPE:
 (I) ORGANEILE:
 W--> 121 (v) FRAGMENT TYPE:
 (1) ORIGINAL SOURCE:
 (2) ORGANISM:
 (3) STRAIN:
 (4) INDIVIDUAL IDENTIFIER:
 (5) DEVELOPMENTAL STAGE:
 (6) HOST TYPE:
 (7) TISSUE TYPE:
 (8) CELL TYPE:
 (9) CELL LINE:
 (I) ORGANEILE:
 C--> 131 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
 C--> 133 N L G E H
 E--> 134

same eno

same

